

A

NotI

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ATG	AAG	CGA	GGA	TGG	AAA	AAG	AAT	TTC	ATA	GCC	GTC	TCA	GCA	GCC	AAC	CGC	TTT	AAG	AAA	ATC	TCA	TCC	TCC
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CBP tag

Gly/Ser-linker

GGG	GCA	CTT
G	A	L

GGA	AGC	GGT	AGC	GGT	ACC
G	S	G	S	G	S

SBP tag (SB19C4)

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M	D	E	K	T	T	G	M	R	G	G	H	V	E	G	L	A

GGC	GAG	CTG	GAG	CTG	CGG	GCC	AGG	CTG	GAG	CAC	CCT	CAG	GGC	CAG	CGG	GAG	CCC
G	E	L	Q	L	R	A	R	L	E	H	P	Q	G	Q	R	E	P

TCC	GGC	GGC	TGC	AAG	CTG	GGC
S	G	G	C	K	L	G

* GCCCGGGCGGATCC

tag length: 78AA

B

XhoI

G/S-linker

CTC	GAG	GGA	AGC	GGT	AGC	GGT	ACC
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SBP-tag

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SBP-tag

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TCC	GGC	GGC	TGC	AAG	CTG	GGC	TCC
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CBP-tag

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Apal

TAG GGCCCGAC

Tag length: 79 AA

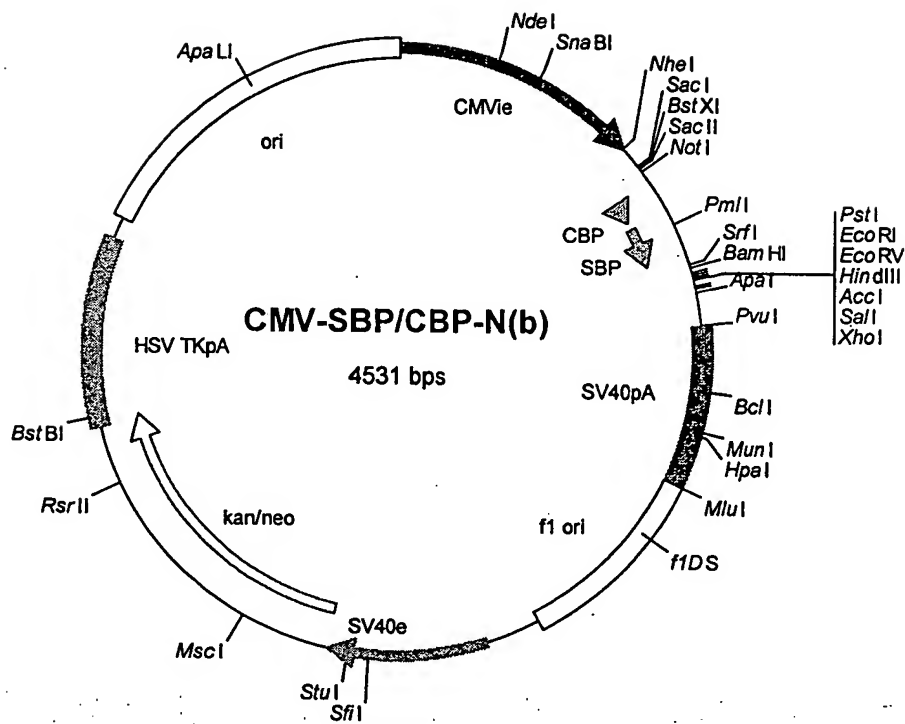
Fig. 1

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SB2	2	<u>MDEKTHCF</u> HPGDHLVRLVEELQALAEGLQGGQPHRLPRRPHHLQLLIDAEHPQAGLRERAHQVDRILLQHHPPQGDRLAQPDHPLLELVWRLLPPS
SB3	4	<u>MTRRPT</u> ASSSSCVRHLLLRQGEHGHQALSDRDKARHVLVEGDVEVLGGDLRLARHREALHPQAGLVHLPGLHGGDLGGHLRLVLEAHPQCDRLGLAVHHH
SB4	1	<u>MDEKTHWGI</u> STWRGEPILLHPQAGRLPLDRRRARHRRILGAEPGGVDHGLRLLELDHRLPLVDHHPQGPQLRGDLPQVVPVLRHARHVLGGLAAATLT
SB5	3	<u>MDEKTHWV</u> NYHPQGDLLVRGHGHVREALHDOGLHQDLQDGLLVGPPPEVVRALRGVILGGLRLVPLDHPQGEALDQARQPQHLLHHRALPPALVWRLLPPS
SB6	1	<u>MDEKTHWLN</u> NPEELARLDGLREGEDHPLVLRHHPQGDGLLDQPLGRHRAIDGVEVREGDRPLDQGGEDIGALVDDDDGEVLDGLVHVGVHVDPLVCGCHHH
SB7	1	<u>MDEKTHW</u> FTLNSPPTHMSAVNGKIDCSFNHNLSTNHWSLSSGHPDGLDDQLEHPQGDALVGRDDGVQALRLLEGQHQRRLAQRRADHRQLVWRLLPPS
SB8	1	<u>MDEKTHCTI</u> ELNPSFTHWKLHHPQGDALLDDGVRPHHPLADEGGGLDQGLGHRGVVAERLARDDPEVLGELVERHGLVPLRHGGHRAEPVLRLLPPS
SB9	1	<u>MDEKTHC</u> NTGLYDGAACFNEALNDVPLVEGRHDLVEGLLERHPQGDPLVAHQVHHPILGRGERHREALVQOEHQPHRLQPVVDLGRRLVWRLLPPS
SB10	1	<u>MDEKTHW</u> HERAQELVGGILLHHPQRLILEPRGPRPLRGVHERGHQPPQLAGRVEEADGGILLRDGGCELEPLVREGEDHLEPLDDELDAGPRGLVWRLLPHHH
SB11	1	<u>MDEKTHW</u> HERVHHLADGLEQHPQGRPLVERHRQVPRGLVRLQHEGLPIEHPAGVHVIRLHQDDRDVDGLVDGHRDVRGLEREVDGPHRLVWRLLPPS
SB12	4	<u>MDKDP</u> LEELSELRLVHHPQGGLLPLRGQVGHDAERLGAEVDDLRGLLDEPQRAVAGLHHVPHRVGQRLVHVRVRELDGLDQDDLRQLVWRLLPPS
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SB14	1	<u>MDEKTHRT</u> LSVLSFNDWLGGTKACWRLVGLHGRHPQGLVREHVDVPLAEVQVQVVGGLADGVEQPGGILLHRAQRVDHPLPDHAGQVLRVWRLLPPS
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SB16	1	<u>MDEKTHW</u> VLQLHPQGDRLGPRHGGDVLRLVCGGEGVLEGLDGRPRRRHRLPREDEHVRALVDQVRLAERLVEEVGGVEALRHGLPQDEPRSGGCHHH
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SB18	1	<u>MDEKTHC</u> AVNVNVELTWHCHRVHHLQPLDPEHPQGDHLRLPLGHALVDPLVQGVVEVVRPLQDVGVQRVALVEQVAVVEGLDHEAGAHGALVWRLLPPS
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SB20	1	<u>MNEKTHC</u> KLNFKVNIADWLAEFHHGGGGLGRRDGVVQLVDGVQVERLDRDPGLGDLRLLELHHRDLRLGGEHLRLRDHPLSPDDHLLVVGGLVWRLLPPS

The "No." indicates the number of times each sequence was observed. The HPQ sequence is in bold type. Defined sequences at the termini are underlined. The six C-terminal residues are not shown.

Fig. 2

Fig. 3a-1



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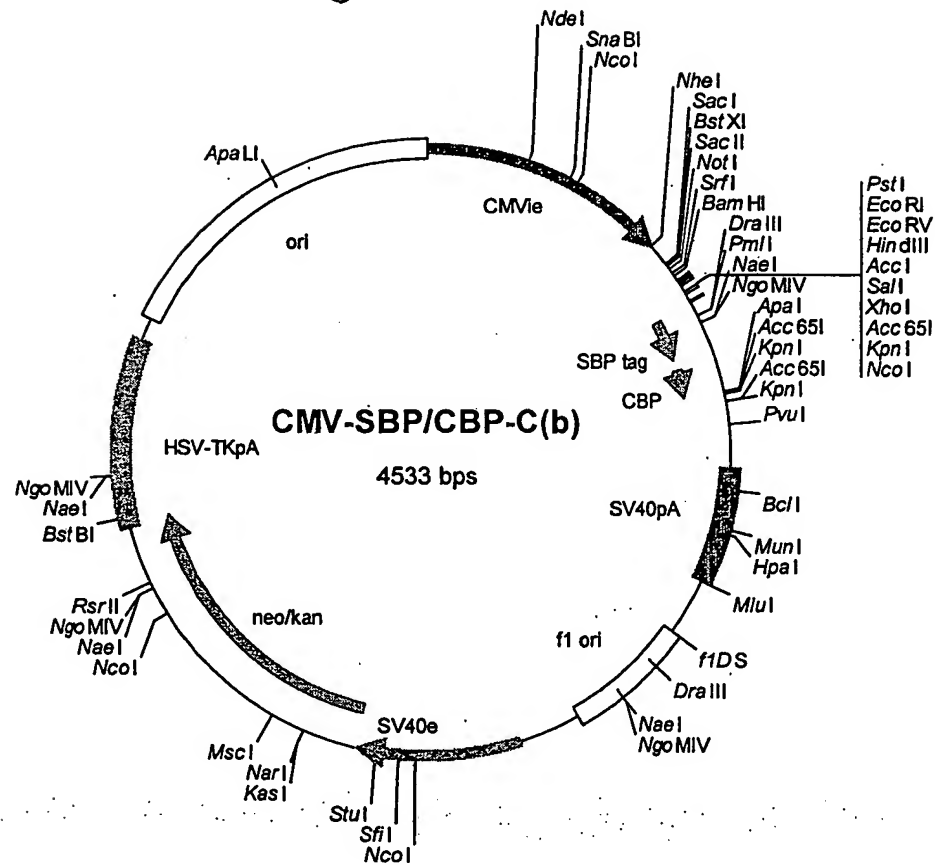
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Fig. 3a-2

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Fig. 3a-3

Fig. 3b-1



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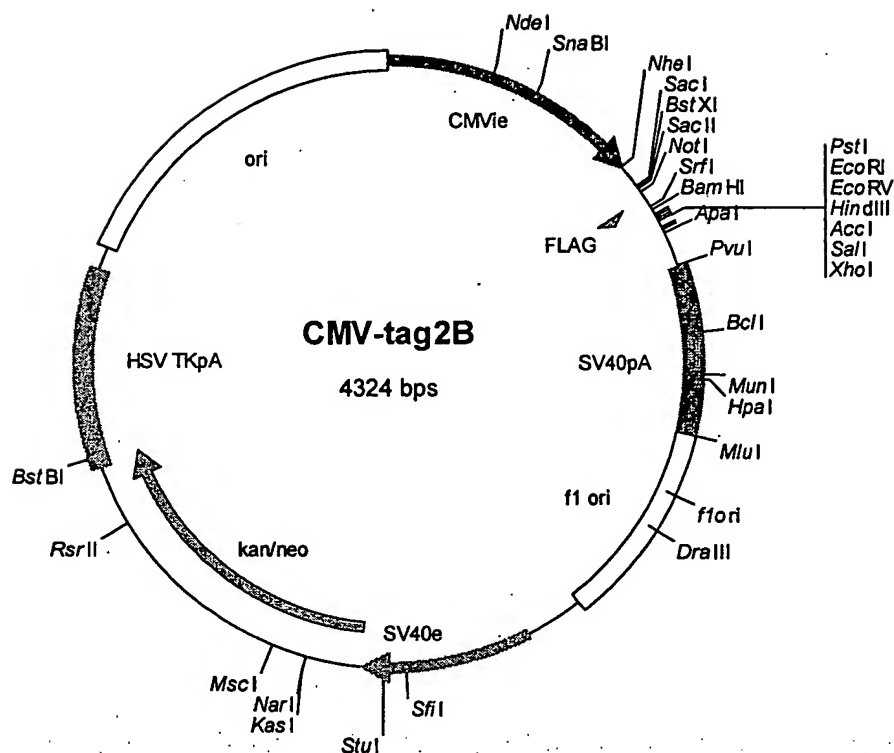
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Fig. 3b-2

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Fig. 3b-3

Fig. 4a-1



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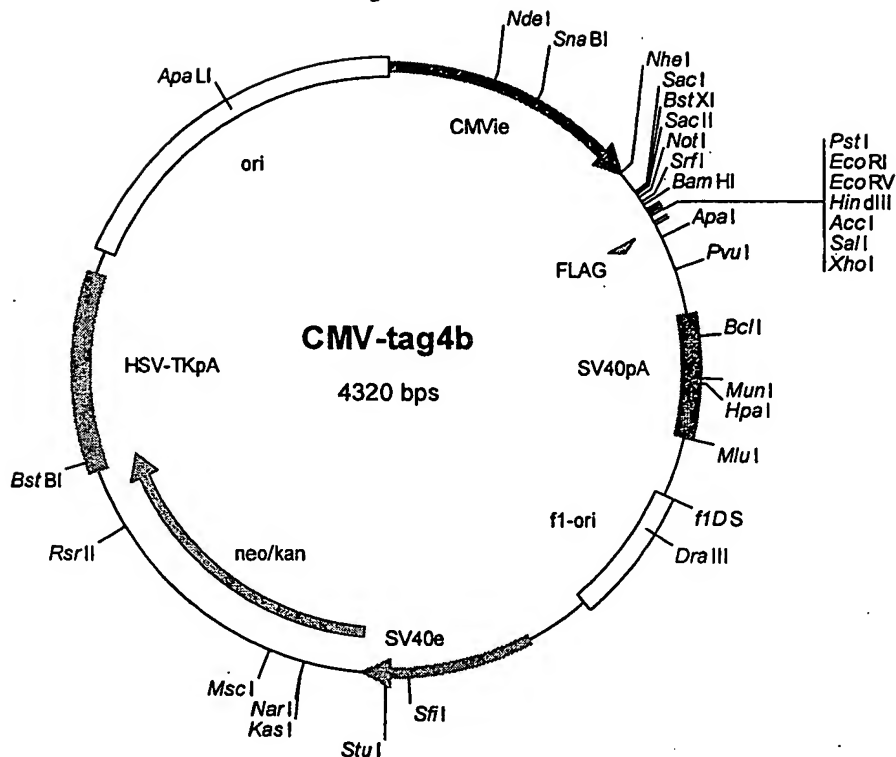
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Fig. 4a-2

Fig. 4b-1



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Fig. 4b-2

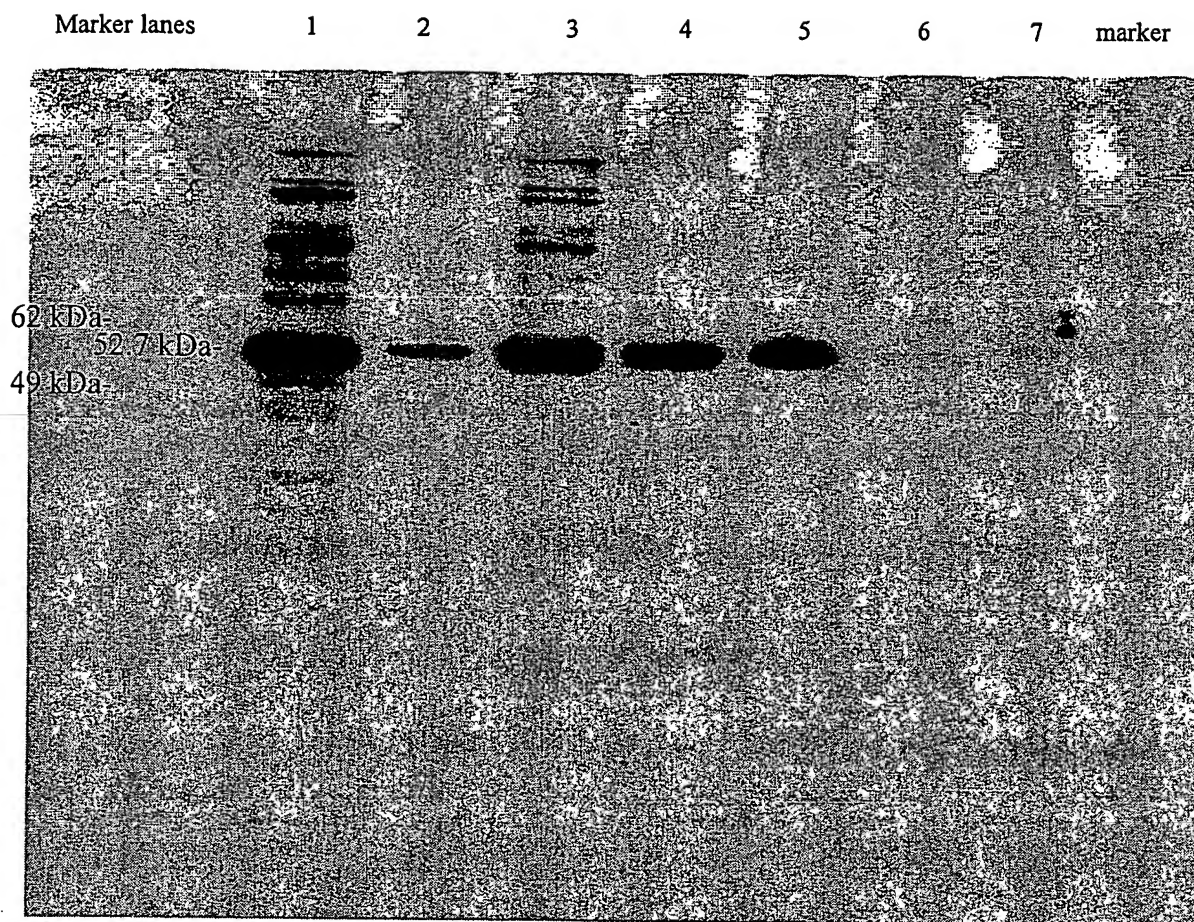


FIGURE 5

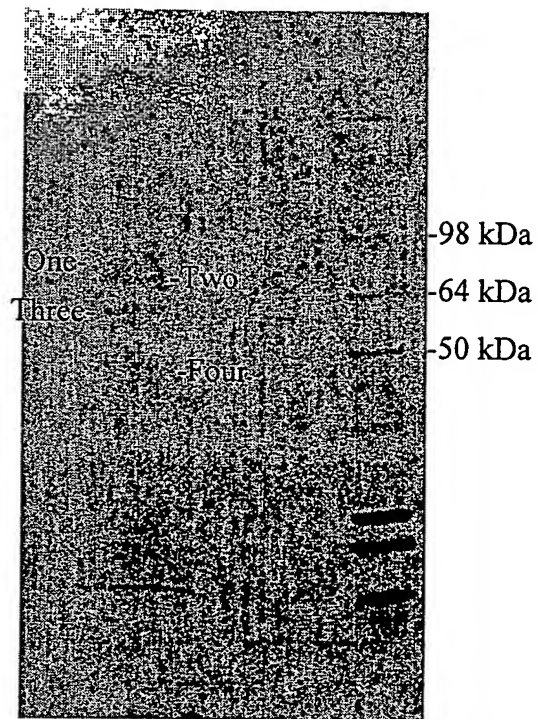


FIGURE 6